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SEQUENCE LISTING

<110> Petrini, John H.J.
Morgan, William Franklin
Maser, Richard Scott
Carney, James Patrick

<120> DNA Encoding A DNA Repair Protein

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<140> US 09/067,641

<141> 1998-04-27

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<213> Homo sapiens

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<212> PRT

<213> Homo sapiens

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			20					25					30		

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 Thr Ala Asn Phe Ser Val Thr Asn Leu Ser Gln Thr Asp Glu Ile Pro
 50 55 60
 Val Leu Thr Leu Lys Asp Asn Ser Lys Tyr Gly Thr Phe Val Asn Glu
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 Glu Lys Met Gln Asn Gly Phe Ser Arg Thr Leu Lys Ser Gly Asp Gly
 85 90 95
 Ile Thr Phe Gly Val Phe Gly Ser Lys Phe Arg Ile Glu Tyr Glu Pro
 100 105 110
 Leu Val Ala Cys Ser Ser Cys Leu Asp Val Ser Gly Lys Thr Ala Leu
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 Asn Gln Ala Ile Leu Gln Leu Gly Gly Phe Thr Val Asn Asn Trp Thr
 130 135 140
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 145 150 155 160
 Thr Ile Cys Ala Leu Ile Cys Gly Arg Pro Ile Val Lys Pro Glu Tyr
 165 170 175
 Phe Thr Glu Phe Leu Lys Ala Val Gln Ser Lys Lys Gln Pro Pro Gln
 180 185 190
 Ile Glu Ser Phe Tyr Pro Pro Leu Asp Glu Pro Ser Ile Gly Ser Lys
 195 200 205
 Asn Val Asp Leu Ser Gly Arg Gln Glu Arg Lys Gln Ile Phe Lys Gly
 210 215 220
 Lys Thr Phe Ile Phe Leu Asn Ala Lys Gln His Lys Lys Leu Ser Ser
 225 230 235 240
 Ala Val Val Phe Gly Gly Gly Glu Ala Arg Leu Ile Thr Glu Glu Asn
 245 250 255
 Glu Glu Glu His Asn Phe Phe Leu Ala Pro Gly Thr Cys Val Val Asp
 260 265 270
 Thr Gly Ile Thr Asn Ser Gln Thr Leu Ile Pro Asp Cys Gln Lys Lys
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 Asn Tyr Cys Asp Pro Gln Gly His Pro Ser Thr Gly Leu Lys Thr Thr

325 330 335
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 Ser Glu Gln Ala Asp Thr Trp Asp Leu Ser Glu Arg Pro Lys Glu Ile
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 Pro Thr Val Lys Glu Ser Cys Lys Thr Ser Ser Asn Asn Asn Ser Met
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 Pro Thr Lys Leu Pro Ser Ile Asn Lys Ser Lys Asp Arg Ala Ser Gln
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 Lys Arg Glu Arg Asp Glu Glu Asn Gln Glu Met Ser Ser Cys Lys Ser
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 Thr Pro Ser Leu Trp Lys Asn Lys Glu Gln His Leu Ser Glu Asn Glu
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 Pro Val Asp Thr Asn Ser Asp Asn Asn Leu Phe Thr Asp Thr Asp Leu
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 Lys Ser Ile Val Lys Asn Ser Ala Ser Lys Ser His Ala Ala Glu Lys
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 Leu Arg Ser Asn Lys Lys Arg Glu Met Asp Asp Val Ala Ile Glu Asp
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 Asp Val Lys Val Gln Lys Gln Glu Glu Asp Val Asn Val Arg Lys Arg
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 Thr Glu Phe Arg Ser Leu Val Ile Lys Asn Ser Thr Ser Arg Asn Pro
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 Lys Lys Val Thr Tyr Pro Gly Ala Gly Lys Leu Pro His Ile Ile Gly
 690 695 700
 Gly Ser Asp Leu Ile Ala His His Ala Arg Lys Asn Thr Glu Leu Glu
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 Glu Trp Leu Arg Gln Glu Met Glu Val Gln Asn Gln His Ala Lys Glu
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<213> Homo sapiens

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35 40 45
Glu Asn Glu Val Asp Phe Ile Leu Leu Gly Gly Asp Leu Phe His Glu
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Asn Lys Pro Ser Arg Lys Thr Leu His Thr Cys Leu Glu Leu Leu Arg
65 70 75 80
Lys Tyr Cys Met Gly Asp Arg Pro Val Gln Phe Glu Ile Leu Ser Asp
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Gln Ser Val Asn Phe Gly Phe Ser Lys Phe Pro Trp Val Asn Tyr Gln
100 105 110
Asp Gly Asn Leu Asn Ile Ser Ile Pro Val Phe Ser Ile His Gly Asn
115 120 125
His Asp Asp Pro Thr Gly Ala Asp Ala Leu Cys Ala Leu Asp Ile Leu
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Ser Cys Ala Gly Phe Val Asn His Phe Gly Arg Ser Met Ser Val Glu
145 150 155 160
Lys Ile Asp Ile Ser Pro Val Leu Leu Gln Lys Gly Ser Thr Lys Ile
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Ala Leu Tyr Gly Leu Gly Ser Ile Pro Asp Glu Arg Leu Tyr Arg Met
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Phe Val Asn Lys Lys Val Thr Met Leu Arg Pro Lys Glu Asp Glu Asn
195 200 205
Ser Trp Phe Asn Leu Phe Val Ile His Gln Asn Arg Ser Lys His Gly
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Ser Thr Asn Phe Ile Pro Glu Gln Phe Leu Asp Asp Phe Ile Asp Leu
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Val Ile Trp Gly His Glu His Glu Cys Lys Ile Ala Pro Thr Lys Asn
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Glu Gln Gln Leu Phe Tyr Ile Ser Gln Pro Gly Ser Ser Val Val Thr
260 265 270

Ser Leu Ser Pro Gly Glu Ala Val Lys Lys His Val Gly Leu Leu Arg
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 Ile Lys Gly Arg Lys Met Asn Met His Lys Ile Pro Leu His Thr Val
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 Arg Gln Phe Phe Met Glu Asp Ile Val Leu Ala Asn His Pro Asp Ile
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 Phe Asn Pro Asp Asn Pro Lys Val Thr Gln Ala Ile Gln Ser Phe Cys
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 Gly Asn Ser His Gln Pro Glu Lys Pro Leu Val Arg Leu Arg Val Asp
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 485 490 495
 Ile Asp Glu Glu Val Arg Arg Phe Arg Glu Thr Arg Gln Lys Asn Thr
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Thr Ser Ser Ser Lys Ile Met Ser Gln Ser Gln Val Ser Lys Gly Val		
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 Glu Asn Glu Val Asp Phe Ile Leu Leu Gly Gly Asp Leu Phe His Glu
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 Asn Lys Pro Ser Arg Lys Thr Leu His Thr Cys Leu Glu Leu Leu Arg
 65 70 75 80

Lys Tyr Cys Met Gly Asp Arg Pro Val Gln Phe Glu Ile Leu Ser Asp
 85 90 95
 Gln Ser Val Asn Phe Gly Phe Ser Lys Phe Pro Trp Val Asn Tyr Gln
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 Asp Gly Asn Leu Asn Ile Ser Ile Pro Val Phe Ser Ile His Gly Asn
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 His Asp Asp Pro Thr Gly Ala Asp Ala Leu Cys Ala Leu Asp Ile Leu
 130 135 140
 Ser Cys Ala Gly Phe Val Asn His Phe Gly Arg Ser Met Ser Val Glu
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 Lys Ile Asp Ile Ser Pro Val Leu Leu Gln Lys Gly Ser Thr Lys Ile
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 Ala Leu Tyr Gly Leu Gly Ser Ile Pro Asp Glu Arg Leu Tyr Arg Met
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 Ser Trp Phe Asn Leu Phe Val Ile His Gln Asn Arg Ser Lys His Gly
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 Ser Thr Asn Phe Ile Pro Glu Gln Phe Leu Asp Asp Phe Ile Asp Leu
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 Val Ile Trp Gly His Glu His Glu Cys Lys Ile Ala Pro Thr Lys Asn
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 Arg Gln Phe Phe Met Glu Asp Ile Val Leu Ala Asn His Pro Asp Ile
 305 310 315 320
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 325 330 335
 Leu Glu Lys Ile Glu Glu Met Leu Glu Asn Ala Glu Arg Glu Arg Leu
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 Gly Asn Ser His Gln Pro Glu Lys Pro Leu Val Arg Leu Arg Val Asp
 355 360 365
 Tyr Ser Gly Gly Phe Glu Pro Phe Ser Val Leu Arg Phe Ser Gln Lys

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 Glu Asp Ile Phe Pro Thr Thr Ser Lys Thr Asp Gln Arg Trp Ser Ser
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Phe Met Thr Thr Lys

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